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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

: Patent Application of

Eric Garver et al.

Conf. No.:

: 5387

Appln. No.

: 09/932,300

Filed

For

: August 20, 2001

: Methods of Inhibiting Alcohol Consumption

COPY OF PAPERS ORIGINALLY FILED

Group Art Unit: Not

Yet Assigned

Attorney Docket No.

9855-3U2

SECOND PRELIMINARY AMENDMENT

Prior to examination on the merits of the application referenced above, please amend the application as follows.

In the Specification:

Please delete paragraphs [0048] and [0049] and substitute the following paragraphs [0048] and [0049] in place thereof. For the Examiner's convenience, the Applicants have enclosed a marked-up version of the replacement specification paragraphs, wherein deleted text is struck-through and added text is underlined.

Figure 11, comprising Figures 11A and 11B, is the sequence (SEQ ID NO: 108) [0048] disclosed in Farres et al. (1989, Eur. J. Biochem. 180:67-74) for the mRNA encoding rat mitochondrial aldehyde dehydrogenase 2 (GENBANK® accession no. NM_032416). The underlined residues are residues 241-261 and have the sequence SEQ ID NO: 107.

Figure 12, comprising Figures 12A and 12B, is the sequence (SEQ ID NO: 110) for 100491 the mRNA encoding human mitochondrial aldehyde dehydrogenase 2 (GENBANK® accession no. NM 000690). The underlined residues are residues 237-257, have the sequence SEQ ID NO: 109, and exhibit high homology with SEQ ID NO: 107.

Remarks

No new matter has been added. The replacement specification paragraphs simply conform the figure legends with the formalized figures.

Respectfully submitted,

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2 January 2002 (Date)

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Marked-Up Copy of Replacement Specification Paragraphs

[0048] Figure 11, comprising Figures 11A and 11B, is the sequence (SEQ ID NO: 108) disclosed in Farres et al. (1989, Eur. J. Biochem. 180:67-74) for the mRNA encoding rat mitochondrial aldehyde dehydrogenase 2 (GENBANK® accession no. NM_032416). The underlined residues are residues 241-261 and have the sequence SEQ ID NO: 107.

[0049] Figure 12, comprising Figures 12A and 12B, is the sequence (SEQ ID NO: 110) for the mRNA encoding human mitochondrial aldehyde dehydrogenase 2 (GENBANK® accession no. NM_000690). The underlined residues are residues 237-257, have the sequence SEQ ID NO: 109, and exhibit high homology with SEQ ID NO: 107.